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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*

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5: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*

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APPLICANT: Murison, Greg FILE OF INVENTION: Fibroblast Growth Factor Rec FILE REFERENCE: 11000.1037C5 CURRENT APPLICATION NUMBER: US/10/613,413A CURRENT FILING DATE: 2003-07-03 PRIOR APPLICATION NUMBER: U.S. 09/823,038 PRIOR FILING DATE: 2001-03-28 PRIOR APPLICATION NUMBER: U.S. 09/823,038 PRIOR APPLICATION NUMBER: U.S. 09/383,586 PRIOR APPLICATION NUMBER: U.S. 09/276,268 PRIOR APPLICATION NUMBER: U.S. 09/276,268 PRIOR APPLICATION NUMBER: PCT/NZ00/00015 PRIOR APPLICATION NUMBER: PCT/NZ00/00015 PRIOR APPLICATION NUMBER: U.S. 60/221,216 PRIOR APPLICATION NUMBER: U.S. 60/221,216 PRIOR APPLICATION NUMBER: U.S. 10/157,444 PRIOR APPLICATION NUMBER: PCT/NZ03/00105 PRIOR APPLICATION NUMBER: PCT/NZ03/00105 PRIOR FILING DATE: 2000-05-28 PRIOR FILING DATE: 2003-05-27 NUMBER OF SEQ ID NOS: 145 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 49 LENGTH: 357 TYDE: DRT Sequence 49, Applicate Publication No. US20 GENERAL INFORMATION: Query Matc Best Local Matches 3 10-613-413A-APPLICANT: Sle LENGTH: 357 TYPE: PRT ORGANISM: Human Match 11 S: , in Application US/10613413A 10. US20040058849A1 99.4%; ilarity 99.7%; Conservative eman, Matthew Score 1707; DB 15; Pred. No. 4.1e-108; 0; Mismatches 1; Receptors and Methods Length Indels 357; 0 Gaps for Their 0; Use

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APPLICANT: Saris, Christiaan M.
APPLICANT: Sharon, Mu X.
APPLICANT: Xia, Min
APPLICANT: Boone, Thomas Charles
APPLICANT: Covey, Todd
TITLE OF INVENTION: Fibroblast Growth Fac
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 99-513-A
CURRENT APPLICATION NUMBER: US/09/815,100
CURRENT FILING DATE: 2001-03-22
PRIOR APPLICATION NUMBER: 60/191,379
PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 5
LENGTH: 472
TYPE: PRT
ORGANISM: Homo sapiens
US-09-815-108-5
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US-09-815-108-5
; Sequence 5, Application US/09815108
; Patent No. US20020009776A1
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Best Local Similarity 99.7%;
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RESULT 4
US-09-758-386-2
; Sequence 2, Application US/09758386
; Patent No. US20010016335A1
; GENERAL INFORMATION:
; APPLICANT: Human Genome Sciences, In
; TITLE OF INVENTION: Fibroblast Growth

FILE REFERENCE:
CURRENT APPLICAT
CURRENT FILING I
PRIOR APPLICATION
PRIOR FILING DATE

ATION NUMBER: US/09/758,386 DATE: 2001-01-12 ION NUMBER: 09/293,182

1999-04-16

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US-10-229-584-5
; Sequence 5, Application; Publication No. US200; GENERAL INFORMATION:
; APPLICANT: Saris, Ch
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Best Local
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TYPE: PRT
ORGANISM:
-10-229-584-
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TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 99-513-F
CURRENT APPLICATION NUMBER: US/10/229,584
CURRENT FILING DATE: 2002-08-28
PRIOR APPLICATION NUMBER: 09/815,108
PRIOR FILING DATE: 2001-03-22
PRIOR APPLICATION NUMBER: 60/191,379
PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 22
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APPLICANT: Xia, Min
APPLICANT: Boone, Thomas Charles
APPLICANT: Covey, Todd
TITLE OF INVENTION: Fibroblast Growth Factor
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 99-513-A
CURRENT APPLICATION NUMBER: US/09/815,108
CURRENT FILING DATE: 2001-03-22
PRIOR APPLICATION NUMBER: 60/191,379
PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn Ver. 2.0
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ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description
OTHER INFORMATION: FGFR-L amino
OTHER INFORMATION: 1-472 of SEC
OTHER INFORMATION: GenBank acce
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Matches 323
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Sequence 8,
Patent No.
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TYPE: PRT
ORGANISM: Homo
-09-758-386-2
                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/09815108 Patent No. US20020009776A1 GENERAL INFORMATION:
                                  Query Match

Best Local Similarity 99.7%;

Matches 323; Conservative
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Similarity 99.7%;
23; Conservative
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MTPSPLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL
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                                                                                               Description of Artificial Sequence: virtual hurFGFR-L amino acid sequence comprising residues 1-472 of SEQ ID NO: 5 and residues 473-504 of GenBank accession no. AJ277437
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                                    Score 1707; DB 9; Pred. No. 5.9e-108; 0; Mismatches 1;
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APPLICANT: Saris, Christiaan M.

APPLICANT: Sharon, Mu X.

APPLICANT: Sharon, Mu X.

APPLICANT: Sharon, Mu X.

APPLICANT: Boone, Thomas Charles

APPLICANT: Covey, Todd

ITITLE OF INVENTION: Fibroblast Growth Factor Receptor-Like Molecules and

ITITLE OF INVENTION: Uses Thereof

FILE REFERENCE: 99-513-A

CURRENT APPLICATION NUMBER: US/09/815,108

CURRENT APPLICATION NUMBER: 001-03-22

PRIOR APPLICATION NUMBER: 60/191,379

PRIOR APPLICATION NUMBER: 60/191,379

PRIOR FILING DATE: 2000-03-22

NUMBER OF SEQ ID NOS: 22

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 15

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RESULT 7
US-09-815-108-17
; Sequence 17, Application US/0; Patent No. US20020009776A1
· GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Saris, Christiaan M.
APPLICANT: Sharon, Mu X.
APPLICANT: Sharon, Mu X.
APPLICANT: Xia, Min
APPLICANT: Boone, Thomas Charles
APPLICANT: Covey, Todd
TITLE OF INVENTION: Fibroblast Growth Fa
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 99-513-A
CURRENT APPLICATION NUMBER: US/09/815,10
                                                                                                                            RESULT 8
US-09-815-108-19
Sequence 19, Application US/09815108
Patent No. US20020009776A1
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APPLICANT: Xia, Min
APPLICANT: Boone, Thomas Charles
APPLICANT: Covey, Todd
TITLE OF INVENTION: Fibroblast Growth Fa
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 99-513-A
CURRENT APPLICATION NUMBER: US/09/815,10
CURRENT FILING DATE: 2001-03-22
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Best Local S
Matches 323
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SEQ ID NO 17
LENGTH: 504
TYPE: PRT
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PRIOR FILING DATE: 2000-03-22
NUMBER OF SEQ ID NOS: 22
SOFTWARE: Patertt
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9-815-108-17
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NUMBER OF SEQ ID NOS
SOFTWARE: PatentIn V
SEQ ID NO 19
LENGTH: 504
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APPLICANT:
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APPLICANT:
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: William I.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C63
CURRENT APPLICATION NUMBER: US/09/989,722
CURRENT FILING DATE: 2001-11-19
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Gerritsen, Mary E.
Goddard, Audrey
Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Kljavin, Ivar J.
Napier, Mary A.
Pan, James
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DATE: 2000-03-22

2 ID NOS: 22

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art, Timothy A.
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PRIOR APPLICATION NUMBER: 60/049787
PRIOR PILING DATE: 1997-10-17
PRIOR PRIOR PRILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR PRILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/065311
PRIOR PILING DATE: 1997-11-12
PRIOR PILING DATE: 1997-11-12
PRIOR PILING DATE: 1997-11-24
PRIOR PILING DATE: 1998-03-20
PRIOR PILING DATE: 1998-03-20
PRIOR PILING DATE: 1998-03-20
PRIOR PILING DATE: 1998-05-07
PRIOR PILING DATE: 1998-06-02
PRIOR PILING DATE: 1998-06-03
PRIOR PILING DATE: 1998-06-04
PRIOR PILING DATE: 1998-06-05
PRIOR PILING DATE: 1998-06-10
PRIO

PRIOR FILING DATE: 1998-06-12
PRIOR FILING DATE: 1998-06-16
PRIOR APPLICATION NUMBER: 60/089512
PRIOR APPLICATION NUMBER: 60/089514
PRIOR APPLICATION NUMBER: 60/089514
PRIOR APPLICATION NUMBER: 60/089514
PRIOR APPLICATION NUMBER: 60/089518
PRIOR APPLICATION NUMBER: 60/089538
PRIOR APPLICATION NUMBER: 60/089598
PRIOR APPLICATION NUMBER: 60/089599
PRIOR FILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089600
PRIOR PHILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089600
PRIOR PHILING DATE: 1998-06-17
PRIOR APPLICATION NUMBER: 60/089907
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RESULT 10
US-09-989-723-119
; Sequence 119, Application US/09989723
; Patent No. US20020072092A1
; GENERAL INFORMATION:
   APPLICANT: Ashkenazi, Avi J.
   APPLICANT: Baker, Kevin P.
   APPLICANT: Botstein, David
   APPLICANT: Botstein, David
   APPLICANT: Eaton, Dan L.
   APPLICANT: Ferrara, Napoleone
   APPLICANT: Gerber, Hanspeter
   APPLICANT: Geritsen, Mary E.
   APPLICANT: Goddard, Audrey
   APPLICANT: Goddard, Audrey
   APPLICANT: Godowski, Paul J.
   APPLICANT: Grimaldi, J. Christopher
   APPLICANT: Gurney, Austin L.
   APPLICANT: Kljavin, Ivar J.
   APPLICANT: Napier, Mary A.
   APPLICANT: Pan, James
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R APPLICATION NUMBER: 60/091633
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APPLICANT: Brow, Magnett Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Stewart, Timothy A.
APPLICANT: Watenabb, Colin K.
APPLICANT: Watenabb, Colin K.
APPLICANT: Watenabb, Colin K.
APPLICANT: Wood, William P. Mickey
APPLICANT: Wood, William I.
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R APPLICATION NUMBER: 60/090863
R FILING DATE: 1998-06-26
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R FILING DATE: 1998-07-01
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R APPLICATION NUMBER: 60/091519
R APPLICATION NUMBER: 60/091519
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R APPLICATION NUMBER: 60/091626
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RESULT 11
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; Patent No. US200
; PaterAL INFORMAT APPLICANT: APPLICANT: APPLICANT: Ashkenazi, Avi J. Baker, Kevin P. Botstein, David Application US/09989279 0020072496A1 ATION:

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DATE: 1998-06-05
TION NUMBER: 60/088655
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TE: 1998-06-05 ON NUMBER: 60/088212 TE: 1998-06-05

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APPLICANT: Desnoyers, Luc
APPLICANT: Ferrar, Agoleone
APPLICANT: Ferrar, Nagoleone
APPLICANT: Ferrar, Nagoleone
APPLICANT: Fong, Sherman
APPLICANT: Gerber, Hanspeter
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APPLICANT: Gerintteen, Mary E.
APPLICANT: Goddard, Audrey
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APPLICANT: Pan, James
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APPLICANT: Watanabe, Colin K.
APPLICANT: Watanabe, Colon M.
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APPLICANT Baker, Revin P.
APPLICANT Baker, Kevin P.
APPLICANT Belevin, David
APPLICANT Beckein, David
APPLICANT Beckein, David
APPLICANT Beckein, David
APPLICANT Gerber, Hanspeter
APPLICANT Hans, Daniel
APPLICANT Hans, Daniel
APPLICANT Hans, Daniel
APPLICANT Wood, William I.
APPLICANT WOOD, WILLIAMS, DANIEL
APPLICANT WOOD, WILLIAMS, DOOR, DOOR, DOOR, DOOR, DANIEL
APPLICANTON NUMBER: 60/08220
PRIOR PELLING DATE: 1997-10-17
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PRIOR APPLICATION NUMBER: 60/08516
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PRIOR APPLICATION NUMBER: 60/08021
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APPLICANT: Ashkenazi, Avi J.
APPLICANT: Baker, Kevin P.
APPLICANT: Baker, Kevin P.
APPLICANT: Benoyers, Luc
APPLICANT: Benoyers, Luc
APPLICANT: Eaton, Dan L.
APPLICANT: Eaton, Dan L.
APPLICANT: Centon, Dan L.
APPLICANT: Gerritsen, Mary E.
APPLICANT: Godowski, Paul J.
APPLICANT: Godowski, Paul J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mapier, Mary A.
APPLICANT: Napier, Mary A.
APPLICANT: Roy, Margaret Ann
APPLICANT: Roy, Margaret Ann
APPLICANT: Williams, P. Mickey
APPLICANT: Chang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C57
CURRENT ETLING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/062250
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APPLICANT: Ashkenazi, Avi J; APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
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R FILING DATE: 1998-07-02
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R APPLICATION NUMBER: 60/092182
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Gerber, Hanspeter
Gerritsen, Mary E.
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Goddwski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Kljavin, Ivar J.
Napier, Mary A.
Pan, James
Paoni, Nicholas F.
Roy, Margaret Ann
Stewart, Timothy A.
Tumas, Daniel
Watanabe, Colin K.
Williams, P. Mickey
Wood, William I.
Zhang, Zemin
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KW neuroprot
KW antidiabe
KW tuberculo
KW tuberculo
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XX
OS Homo sapi Human; fibroblast growth factor receptor 5; receptor; FGFR5; cytostatic; neuroprotective; antiinflammatory; dermatological; immunosuppressive; antidiabetic; antirheumatic; antiarthritic; tuberculostatic; tuberculostatic; tuberculostatic; nephrotropic; antiarteriosclerotic; vasotropic; osteopathic; gene therapy.

/note= "Tyrosine kinase 217220 /note= "CAAX box" 261324 /note= "Immunoglobulin c 268271 /note= "CAAX box" A1.	/note= "Tyrosine kinase 217220 /note= "CAAX box" Domain	/note= "Tyrosine kinase 217220 /note= "CAAX box" Domain	/note= "Tyrosine kinase 217220 /note= "CAAX box" Domain 261324 /note= "Immunoglobulin c Region 268271	/note= "Tyrosine kinase Region 217220 /note= "CAAX box" Domain 261324 /note= "Immunoglobulin c	/note= "Tyrosine kinase Region 217220 /note= "CAAX box" Domain 261324	/note= "Tyrosine kinase Region 217220 /note= "CAAX box"	/note= "Tyrosine kinase Region 217220	/note= "Tyrosine kinase		Modified-site		/note=	Modified-site 202		Domain 1	/note= "Heparin	Domain 154171	/note	Domain 1		Region 99102	/noi	Region 5154	/note=	Domain 4410	/no	Peptide 124		XX	OS Homo sapiens.
				.271	= "Immunoglobulin domain"		E "CAAA DOX"	. 220	= "Tyrosine kinase phosphorylation site"		horylation site"	"CAMP- and cGMP-dependent protein Kinase		= "Immunoglobulin domain"	. 228	"Heparin binding domain"		= "CAM binding domain"	.171	"CAAX			54	"Immunoglobulin domain"	106	"Signal		ion/Qualifiers		

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is the protein sequence of human fibroblast growth factor receptor 5 (FGFR5). The invention provides murine and human FGFR5 polypeptides and polynucleotides, as well as modulators of FGFR5 gene expression and binding molecules that specifically bind to and agonise of antagonise FGFR5 polypeptide function. Modulators of FGFR5 polypeptide function include antibodies, scFv and Camelidae heavy chain IgG that specifically bind to FGFR5. They can be used to treat diseases associated with elevated osteopontin expression such as cancer (especially breast cancer, hepatocellular carcinoma and colon cancer), multiple sclerosis, systemic lupus erythematosus, diabetes, rheumatoid arthritis, services, tuberculosis, kidney stones, atherosclerosis, vasculitis, nephritis, arthritis, osteoporosis and osteopetrosis.
  Human;
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receptor-like (FGFR-L) polypeptide (I). (I) and the nucleic acid (II) cencoding (I) are useful for treating, preventing or ameliorating a medical condition including haematopoietic disorder, osteogenesis, co osteogenesis imperfecta, Paget's disease, periodontal dieease, cancer, diabetes, obesity and cachexia. (I) is also useful for condentifying a compound which binds to RGFR-L polypeptide, by contacting conjupeptide to the compound, and determining the activity of the polypeptide when bound to the compound. (II) is useful for modulating contacting conjupeptide when bound to the compound. (II) is useful for modulating contacting contacting conjupeptide of a polypeptide in an animal. A transgenic animal comprising (I) contactivity or FGFR-L polypeptide production, by exposing the transgenic animal to the compound and measuring FGFR-L polypeptide or production in the mammal. (II) is useful for mapping the locations of FGFR-L gene and created genes on chromosomes, as hybridisation probes in diagnostic assays to test for the presence of an FGFR-L nucleic acid molecule in for isolating corresponding FGFR-L polypeptide genes. (I) is useful as tools for isolating corresponding FGFR-L polypeptide genes. (I) is useful as conjugent and for cloning FGFR-L polypeptide genes. (I) is useful as conjugent strategy. The present sequence represents the amino acid sequence of human fibroblast growth factor receptor-like protein as described in the invention.
WWW. WAS SEED TO THE TEST OF T
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polynucleotides encoding them. The PRO sequences of the invention were identified based on extracellular domain homology screening. The PRO sequences have homology with proteins including LDL receptors, TIE ligands and various enzymes. The membrane-bound proteins and receptor molecules are useful as pharmaceutical and diagnostic agents. Receptor communoadhesins, for instance, can be used as therapeutic agents to block membrane-bound proteins can also be comployed for screening of potential peptide or small molecule inhibitors confide the relevant receptor/ligand interaction. The PRO encoding sequences care useful as hybridization probes, in chromosome and gene mapping and in the generation of antisense RNA and DNA. PRO nucleic acid sequences will consolidate the preparation of PRO polypeptides, especially by recombinant techniques
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This is the fibroblast growth factor receptor protein, FGFR-5. The FGFR-5 (C diseases or conditions associated with inappropriate FGFR-5 expression and activity. For example, the nucleic acids (and vectors containing them) and the FGFR-5 polypeptide may be used to treat disorders (c.g. leukemia, colonic cancer, testicular cancer and follicular (lymphomas), autoimmune disorders (e.g. multiple sclerosis and Crohn's disease) viral infections (e.g. herpes viruses), inflammation, graft (versus host disease, acute and chronic graft rejection, ischemic injuries and proliferation, diseases associated with defects in wound healing, mucositosis, defects of angiogenesis, immune dysfunction, endocrine dysfunction and insulin secretion disorders. Anti-FGFR-5 antibodies may calso be used as diagnostic agents for detecting the presence of FGFR-5 polypeptides in samples
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07-JUL-1999;
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05-JAN-2000;
                                                                                                                                                                                                     Human; tumour; diagnosis; neoplastic disease; neoplastic cell growth; proliferation; tumourigenesis; identification; cancer; cytostatic; nootropic; neuroprotective; antiinflammatory; immunosuppressive; immunostimulant; antiangiogenic; leukaemia; lymphoid malignancy; neuronal disorder; glial disorder; astrocytal disorder; angiogenic; hypothalamic disorder; glandular disorder; macrophagal disorder; epithelial disorder; stromal disorder; blastocoelic disorder; inflammatory disorder; immunologic disorder.
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                                                              99WO-US005028.

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                                Baker
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RESULT 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes an isolated antibody that binds to one of the human pro proteins designated pro212, pro290, pro341, pro535, pro619, pro717, pro809, pro830, pro848, pro943, pro1005, pro1009, pro1187, pro1181, pro1187, pro1181, pro1187, pro1181, pro1187, pro1181, pro23, pro39, pro834, pro11710, pro11710, pro2094, pro1187, pro1181, pro23, pro39, pro834, pro1317, pro11710, pro2094, pro2187, pro1187, pro1181, pro2198. Pro antagonists can be used to inhibit tumour cell growth. The pro2198. Pro antagonists can be useful in the treatment, diagnosis and prevention of cancer. The antibodies and other anti-tumour compounds maybe used to treat various conditions, including those characterised by overexpression and/or activation of the amplified Pro genes. Exemplary compounds include benign or malignant tumours (e.g., renal, liver, kidney, bladder, breast, gastric, ovarian, colorectal, prostate, pancreatic, lung, vulva, thyroid, hepatic carcinomas, sarcomas, glioblastomas, and various head and neck tumours), leukaemias and lymphoid malignancies, other disorders such as neuronal, glial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, immunologic disorders, and other glandular, macrophagal, epithelial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, astrocytal, hypothalamic and other glandular, macrophagal, epithelial, astrocytal, and sarcomas, and inflammatory, angiogenic and hybridisation probes used in the isolation of the human pro sequences.

Caracant invention and protein sequences given in the exemplification of the
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Membrane associated protein; secreted INTERCEPT 340; MANGO 003; MANGO 347;
                                           Human MANGO
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                                                                                                                                                                                                                                                                                 ent invention provides the protein and coding sequences for a f membrane associated and secreted proteins from human, mouse and proteins are designated INTERCEPT 340, MANGO 003, MANGO 347, 2, TANGO 295, TANGO 254 and TANGO 378. The proteins are all in signal transduction and the sequences can be used in the cof cardiovascular, renal, hepatic, neural, neoplastic, skeletal atopoietic disorders
OKE
            QKFV
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VLPTGDVWSRPDGSYLNKLL 324
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         VLPTGDVWSRPDGSYLNKPL 324
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Pred. No. 1.5e-117;
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20-DEC-1999;
20-DEC-1999;
16-JAN-2000;
11-FEB-2000;
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24-FEB-2000;
24-FEB-2000;
24-FEB-2000;
15-MAR-2000;
15-MAR-2000;
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Ferrara N, Fong S, G
Grimaldi CJ, Gurney I
Roy MA, Stewart TA,
Zhang Z;
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                  The present invention describes human secreted and transmembrane PRO proteins. The PRO proteins have cytostatic activity. The PRO proteins can be used for targeted delivery of bioactive molecules, such as toxins, radiolabels or antibodies, that cause cell death. PRO nucleotide sequences, and their fragments, can be used as hybridisation probes, in chromosomal and gene mapping, and in the generation of anti-sense RNA and DNA. They may also be used to produce transgenic animals which are used to develop and screen therapeutically useful reagents. The PRO nucleotide and protein sequence can be used for tissue typing and in treating ancer. Anti-PRO antibodies can be used in diagnostic assays. AAF44270 to AAF44470 represent PCR primers and hybridisation probes used in the isolation of human PRO sequences. AAF44087 to AAF44269 and AAB65154 to AAB65300 represent human PRO polynucleotide and protein sequences given in the exemplification of the present invention
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99US-0141037P.

99US-0143048P.

99US-0145698P.

99US-0146222P.

99WO-US021547.

99WO-US021547.

99WO-US028313.

99WO-US03095.

99WO-US03095.

99WO-US000219.

2000WO-US000376.

2000WO-US004341.

2000WO-US005004.

2000WO-US005841.

2000WO-US006884.

2000WO-US006884.
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                                                                                                                                                                                                         es used to produce polypeptides used toxins, radiolabels or antibodies, toell death.
                                                                                                                                                                                                                                                                                                                                                       INC
                                                                                                                                                                                     935pp;
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Gerber H, Gerritsen ME, Goddard A,
AL, Kljavin IJ, Napier MA, Pan J,
Tumas D, Watanabe CK, Williams PM,
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mapping;
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tissue typing;
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Godowski
Paoni NF;
Wood WI;
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RESULT 8
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KW PRO
KW PRO
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Best Local S
Matches 323
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18-FEB-2000;
02-MAR-2000;
30-MAR-2000;
28-JUL-2000;
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                            Composition e.g.breast c PRO1250 or F
                                                                               Ashkenazi
Wood WI;
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N-PSDB; ABKI
               Claim
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                                                          11750.
                            for inhibiting neoplastic cancer and ovarian cancer ipro1337 or its agonist.
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2000WO-US004342.
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              2; 101pp; English.
                                                                                                                                                                                                                                                                                                                                      Goddard
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Pred. No. 1.5e-117;
0; Mismatches 1;
                                                                                        Gurney
                                                                                                                                                                                                                                                                                                  504
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                                      cell growth or treating tumors in mammals comprising PRO943,
                                                                                        Napier
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                                                                                        MA,
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                                                                                         Watanabe
                                                                                                                                                                                                                                   cancer;
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RESULT 9
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Best Local S
Matches 323
                                                                                                                                                                                                            Human; IMXP-888; PRO943; fibroblast growth factor receptor; virucide; antibacterial; fungicide; cytostatic; antirheumatic; antiarthritic; antibacterial; immunosuppressive; neuroprotective; antiinflammatory; antidiabetic; dermatological; antiasthmatic; antiallergic; vasotropic; antiatherosclerotic; antipsoriatic; nootropic; infection; cancer; lupus; graft versus host disorder; inflammatory disorder; rheumatoid arthritis; septic shock; multiple sclerosis; adult respiratory distress syndrome; ARDS; pneumonia; diabetes; asthma; allergy; reperfusion injury; eczema; atherosclerosis; cardiovascular disease; psoriasis; fibrosis; sarcoidosis; Alzheimer's disease; cytokine inducer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         neoplastic cell growth or treating tumours in mammals. The method comprises a polypeptide PRO943, PRO1250 or PRO1337 of the invention in admixture with a carrier. The invention also comprises the nucleotide and protein sequences of the three PRO proteins. The composition of the invention is useful for inhibiting neoplastic cell growth and for treating tumours e.g. breast cancer, ovarian cancer, colorectal cancer, lung cancer, central nervous system cancer, melanoma or leukaemia in a mammal. An expression vector containing the nucleotide sequences of the invention is useful for producing the PRO proteins by recombinant techniques, these proteins may then be used to create anti-PRO antibodies which may be used for detection of PRO protein or inhibition of PRO protein activity. The present sequence represents the human PRO943 protein of the invention
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Pred. No. 1.5e-117;
Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel method for activating the immune system in a mammal. The method comprises administering to the mammal an effective amount of a cytokine inducer polypeptide, IMXP-888, or treating an inflammatory disorder in a mammal comprising, administering an effective amount of an IMXP-888 antagonist to the mammal. The method of the invention has virucide, antibacterial, fungicide, cytostatic, antirheumatic, antiarthritic, antibacterial, fungicide, cytostatic, antiartheumatic, antiallergic, vasotropic, antiatherosclerotic, antiasthmatic, antiallergic, vasotropic, antiatherosclerotic, antipsoriatic and nootropic activity. The method is useful for activating the immune system in a mammal having a condition selected from viral, bacterial or fungal infection, cancer, and graft versus host disorders, or for treating an inflammatory disorder in a mammal, preferably human. The method is useful for treating inflammatory disorders such as rheumatoid arthritis, septic shock, multiple sclerosis, adult respiratory distress syndrome (ARDS), pneumonia, diabetes, lupus, asthma and other cardiovascular diseases, eczema, psoriasis, fibrosis and the range of fibrotic disorders, sarcoidosis, Alzheimer's disease, and cancer. This sequence represents member of the IMXP-888 family, PRO943, used in the method of the invention
                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 323
                                                                                                                                                                                                                                                                                                                                                       lung conditions, cardiovascular di fibrotic disorder sequence represen method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Activating immune system in a mammal by administering to the mammal a cytokine inducer polypeptide, IMXP-888, or treating inflammatory disorder in a mammal by administering an IMXP-888 antagonist to the mammal.
                                                                                                                                                                                                                                                                                                                                    Sequence
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376. .396
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11-FEB-2000;
18-FEB-2000;
22-FEB-2000;
24-FEB-2000;
15-MAR-2000;
15-MAR-2000;
17-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; PRO; antiinflammatory; ophthalmological; vasotropic; retinal cell injury; ocular disease; retinitis pigmentosa; macular degeneration; retinal detachment; retinal tear; retinopathy; retinal degenerative disease; macular hole; degenerative myopia; acute retinal necrosis syndrome; traumatic chorioretinopathy; purtscher's retinopathy; oedema; ischaemic condition; retinal vision occlusion; collagen vascular disease; thrombocytopaenic purpura; uveitis; retinal vasculitis; Eales diseas systemic lupus erythematosus; environmental trauma.
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Kljavin I
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8 or PRO1132 polypeption
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  the
                                                                     Fig
                                                                                        survival of retinal cells, or delaying or death, by contacting retinal cells 346, 322, 536, 943, 840, 828, 826, 1068
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99WO-US020944.
99WO-US021090.
99WO-US028313.
99WO-US028301.
2000WO-US0028301.
2000WO-US000376.
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          to promoting the survival of retinal cells, or retinal cell injury or death, by contacting polypeptide such as PRO175, PRO220, PRO216, PRO322, PRO536, PRO943, PRO840, PRO828, PROPERTIES Also included are the nucleic acidal properties.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cell comprising the vector, and anti-PRO antibody. The PRO proteins are useful for promoting survival of retinal cells (retinal neurons such as retinal ganglion cells, displaced retinal ganglion cells, amacrine cells, displaced retinal ganglion cells, amacrine cells, considered amacrine cells, horizontal neurons or bipolar neurons, rod photoreceptors, or supportive cells such as Muller cells or pigment caused by ocular disease (which is or is associated with retinitis retinopathy, retinal degeneration, retinal detachment, retinal tear, contusion, purtscher's retinopathy, oedema, an ischaemic condition, contusion, purtscher's retinopathy, oedema, an ischaemic condition, contusion, purtscher's retinopathy, oedema, an ischaemic condition, contusion branch retinal vision occlusion, collagen vascular disease, thrombocytopaenic purpura, uveitis, retinal vasculitis, occlusion associated with Eales disease or systemic lupus erythematosus), retinal cell injury or death is delayed or prevented by substantially not causing angiogenesis or mitogenesis. The present sequence represents a pRO protein
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cardiac insufadrenal cortion vascular endo endothelial cortinal neuro retinal disor mammalian kiddermatitis; h chondrocyte r ifficiency disorder; cancer; tumour; immune response; ical capillary endothelial growth; c-fos induction; lothelial growth factor inhibition; VEGF inhibition; cell growth inhibitor; T-lymphocytes stimulation; cons cell survival; rod photoreceptor cell survival; order; retinitis pigmentosum; kidney disorder; ldney mesangial cell proliferation; Berger disease; herpetiformis; Crohn's disease; chondrocyte proliferation; redifferentiation; sports injury; arthritis.

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The invention describes an isolated human PRO polypeptide. The PRO C polypeptides are useful in detecting PRO polypeptides in a sample, in C linking a bioactive molecule to a cell expressing a PRO polypeptide, and C in modulating at least one biological activity of a cell expressing a PRO cell expressing a PRO polypeptide. PRO 1312 stimulates hypertrophy of neonatal heart and is thus C useful for treating cardiac insufficiency disorders. PRO1154 and PRO1186 C PRO943, PRO828, PRO826, PRO1068 or PRO535, PRO826, PRO819, PRO1126, PRO943, PRO828, PRO826, PRO1068 or PRO535, PRO826, PRO819, PRO1126, PRO1360 and PRO1387 induce c-fos in endothelial cells, and are thus useful for treating conditions or disorders where angiogenesis would be beneficial, e.g. wound healing and antagonist of this polypeptide are useful for treating cancerous tumours. PRO812 inhibits vascular endothelial growth factor (VEGF) stimulated proliferation of endothelial cells and is thus useful for inhibiting endothelial cell growth in mammals which would be beneficial in inhibiting tumour growth. PRO826, PRO1068, PRO1184, PRO1346 and PRO1375 stimulate proliferation of
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AL, Kljavin IJ, Napier MA, Pan J,
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                                                                                                                                                                                                                                                                      Human; PRO; secreted protein; transmembrane protein; cardiac insufficiency disorders; angiogenesis; wound healing; cancerous tumour; immune response; retinal disorder; sight lo retinitis pigmentosum; age-related macular degeneration; AMD; kidney disorder; Berger disease; nephropathy; dermatitis; her Crohn's disease; sports injury; arthritis.
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                                                                                                                                          The invention discloses isolated PRO secreted/transmembrane polypeptides comprising a sequence without signal peptide and the nucleic acid encoding them. The polypeptides can be used to raise antibodies that specifically bind to the PRO polypeptide, for linking a bioactive molecule to a cell expressing a PRO protein and for modulating at least one biological activity of a cell. The PRO polypeptides or collynucleotides are also useful in gene therapy, in chromosome collypeptides are useful as molecular markers for protein electrophoresis, and the isolated nucleic acids may be used for recombinantly expressing those markers. The PRO polypeptides and nucleic acids may also be used in the isolated nucleic acids may be used for recombinant parents for protein electrophoresis, and the typing. Anti-PRO antibodies are useful in diagnostic assays for PRO, and in affinity purification of PRO from recombinant cell culture or natural sources. The sequences presented in ABUG0478-ABUG0624 are the PRO polymucleotides of the invention. Note: The sequence data for this patent company and also available in electronic format from USPTO at
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Ferrara N,
Grimaldi J
Roy MA, Si
Zhang Z;
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Ashkenazi Ferrara N Grimaldi Roy MA, Zhang Z; N i AJ, Baker N, Fong S, JC, Gurney Stewart TA, KP, I ΑL, (P, Bots Berber H, L, Klja Tumas D, stein D, Desnoyers L, Eaton , Gerritsen ME, Goddard A, avin IJ, Napier MA, Pan J, , Watanabe CK, Williams PM, DL; Godowski Paoni NF; Wood WI; WI;

N-PI SDB 003 A 3-102117/09 ABX64021.

> FFFXXXOOOOOOOOOOOOOOOO S The present invention relates to the isolation of novel human PRO C polypeptides, and the polynucleotide sequences encoding them. The PRO C polypeptides are secreted and transmembrane proteins. The PRO C polypeptides are useful for detecting other PRO polypeptides, for linking bioactive molecules to cells expressing PRO polypeptides, for modulating biological activities of cells expressing PRO polypeptides, and for for identifying agonists or antagonists. The polynucleotide sequences cencoding PRO polypeptides are useful as hybridisation probes, in the preparation of PRO polypeptides, for generating transgenic animals in the preparation of PRO polypeptides, for generating transgenic animals or knockout animals, to construct hybridisation probes for mapping the gene which encodes the PRO polypeptides, for generating transgenic animals identification, as chromosome markers, in gene therapy, for chromosome identification, as chromosome markers, and for generating probes for PCR, Northern analysis, Southern analysis and Western analysis. ABU13860-ABU14006 represent the human PRO polypeptides of the invention. Note: The sequence data for this patent was obtained in electronic format directly from the USPTO web site at seqdata.uspto.gov/psipsDIDEntry.html Query Ma Best Loc Matches Novel secret activity of antagonists Sequence Claim Loca Match $\omega \vdash$. 323 MTP imi ហ Fig 04 ted and transmembrane polypeptide for modulating biological cell expressing the polypeptide, identifying agonists or of polypeptide, and as molecular weight markers. 99.4%; larity 99.7%; Conservative AA; 70; 649pp; English. 0, Score 1707; DB 6; Pred. No. 1.5e-117; 0; Mismatches 1; Length 504; Indels o ; Gaps 60 0

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ALIGNMENTS

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RESULT 1
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; Patent No. 6797271
; GENERAL INFORMATION:
; APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
; APPLICANT: Abernethy, Nevin
; APPLICANT: Murison, Greg
; TITLE OF INVENTION: Compositions Isolat
; TITLE OF INVENTION: and Methods For TI
; FILE REFERENCE: 11000.1037c3
; CURRENT APPLICATION NUMBER: US/09/823,(
CURRENT FILING DATE: 2001-07-09
; NUMBER OF SEQ ID NOS: 61
; SOFTWARE: FastSEQ for Windows Version '
; SEQ ID NO 60
; LENGTH: 373
                                                                                                                                                                                                                                                                                                                             TYPE: PRT
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APPLICANT: Sleeman, Matthew;
APPLICANT: Abernethy, Nevin;
APPLICANT: Onrust, Rene;
APPLICANT: Conrust, Rene;
APPLICANT: Murison, Greg;
TITLE OF INVENTION: Compounds isol;
FILE REFERENCE: 11000.1037c1;
CURRENT APPLICATION NUMBER: US/09;
CURRENT FILING DATE: 1999-08-26;
NUMBER OF SEQ ID NOS: 38;
SOFTWARE: FastSEQ for Windows Vers;
SEQ ID NO 31
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APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Abernethy, Nevin
APPLICANT: Abernethy, Nevin
APPLICANT: Conrust, Rene
APPLICANT: Kumble, Anand
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compositions Isolated From Title OF INVENTION: and Methods For Their Userice Reference: 11000.1037c3
CURRENT APPLICATION NUMBER: US/09/823,038A
CURRENT FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 61
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Best Local Similarity 91.4%;
Matches 296; Conservative
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equence 31, A
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Pred. No. 1.4e-134;
); Mismatches 15;
                                                           Isolated From For Their Use
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RESULT 4
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LENGTH: 529
TYPE: PRT
ORGANISM: Mouse
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SEQ ID NO 33
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Best Local Simil
Matches 273; C
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Best Local S
Matches 296
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                                                                                                                                                                                                                                                                                         Sequence 33, Application US/09383586 Patent No. 6242419
GENERAL INFORMATION:
                                                                                                                          LENGTH: 322
TYPE: PRT
ORGANISM: Human
-09-383-586-33
                                                                                                                                                                                                        APPLICANT: Onrust, Rene
APPLICANT: Kumble, Anand
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compounds isolated from stromal
TITLE OF INVENTION: and methods for their use
FILE REFERENCE: 11000.1037c1
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APPLICANT: Sleeman, Matthew
APPLICANT: Abernethy, Nevin
APPLICANT: Onrust, Rene
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LING DATE: 1999-08-26
SEQ ID NOS: 38
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larity 100.0%;
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Pred. No. 1.4e-134;
9; Mismatches 15;
                                                                                       Score 1458; DB 3; 1
Pred. No. 1.5e-125;
0; Mismatches 0;
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LON:

LICANT: Sleeman, Matthew

LANT: Abernethy, Nevin

LICANT: Onrust, Rene

APPLICANT: Kumble, Anand

APPLICANT: Murison, Greg

TITLE OF INVENTION: Compositions Is.

TITLE OF INVENTION: and Methods For

FILE REFERENCE: 11000.1037c3

CURRENT APPLICATION NUMBER: US/09/823

CURRENT FILING DATE: 2001-07-09

NUMBER OF SEQ ID NOS: 61

SOFTWARE: FastSEQ for Windows Very

SEQ ID NO 33

LENGTH: 322

TYPE: PRT

ORGANISM: Human

3-09-823-038A-33

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est L-

1100.1037c3

CURRENT FILING DATE: 2001-07-09

NUMBER OF SEQ ID NOS: 61

SOFTWARE: FastSEQ for Windows Very

SEQ ID NO 33

LENGTH: 322

TYPE: PRT

ORGANISM: Human

3-09-823-038A-33
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US-09-823-038A-33
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Sequence 32, Application US/09383586; Patent No. 6242419; GENERAL INFORMATION:
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Abernethy, Nevin
APPLICANT: Onrust, Rene
APPLICANT: Kumble, Anand
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compounds isolate
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US-09-383-586-32
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Best Local Similarity 100.0%;
Matches 273; Conservative
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2001-07-09
: 61
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; LENGTH: 439
; TYPE: PRT
; ORGANISM: Mouse
US-09-383-586-32
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; SEQ ID NO 32
; LENGTH: 439
; TYPE: PRT
; ORGANISM: Mouse
US-09-823-038A-32
                                                                                                                                                                                                                                                                                                  ; sequence 32, Application US/09823038A; Patent No. 6797271; GENERAL INFORMATION:
APPLICANT: Strachan, Lorna
APPLICANT: Sleeman, Matthew
APPLICANT: Abernethy, Nevin
APPLICANT: Murison, Greg
TITLE OF INVENTION: Compositions Isolate
TITLE OF INVENTION: and Methods For The
FILE REFERENCE: 11000.1037c3
CURRENT APPLICATION NUMBER: US/09/823,03;
CURRENT FILING DATE: 2001-07-09
NUMBER OF SEQ ID NOS: 61
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US-09-823-038A-32
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Matches 211;
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FILE REFERENCE:
CURRENT APPLICA
CURRENT FILING
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Best Local
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                                                                                         60.3%; nilarity 65.1%; Conservative
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ATION NUMBER: US/09/383,586
DATE: 1999-08-26
                                  PSPLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL
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   LLLGALPSAEAAR
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                                                                                         Score 1035.5; I
Pred. No. 1.3e-8
4; Mismatches
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RESULT 8
US-09-383-630-6
; Sequence 6, Application US/09383630A
; Patent No. 6265632
; GENERAL INFORMATION:
; APPLICANT: Avner Yayon et al.
; APPLICANT: NODE FACTOR RECF
                                                            ; SEQUENCE US-09-383-630-6
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Best Loc
Matches
                                                                                                                                                                    INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                  COUMING SYSTEM: Windows version 2
               Loca
                                                                                                                                            APPLICATION NUMBER: US/09/383,630A
FILING DATE: 26-Aug-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: <Unknown>
FILING DATE: <Unknown>
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Friedmam, Mark M.
REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 1402/2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 972-3-562553
TELEFAX: 972-3-562554
TELEX: <Unknown>
FORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
TENCTH: 801
                              Match
                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: Mark M.
STREET: 2001 Jeffer
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23.8%; 23.1%; 34.4%; 106; Conservative
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                                                                          TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
ENCE DESCRIPTION: SEQ
                                                                                                                       TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Arlington
STATE: Virginia
COUNTRY: United
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                                                                                                                                      801
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2001 Jefferson Davis Highway, Suite 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            on et al.
ANIMAL MODEL FOR FIBROBLAST
FACTOR RECEPTOR ASSOCIATED
CHONDRODYSPLASIA
                                                                                                                                                                                                                                                                                                                                                                                                      for Windows
ASCI file
                                                                          SEQ
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     49
 Score 408.5;
Pred. No. 9.2
9; Mismatches
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NO:
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US-09-383-630-3
; Sequence 3, Application US/u>---
; Sequent No. 6265632
; Patent No. 6265632
; GENERAL INFORMATION:
APPLICANT: Avner Yayon et al.
APPLICANT: Avner Yayon et al.
FACTOR RECEPTOR ASSOCIATED
FACTOR RECEPTOR ASSOCIATED
    US-09-383-
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                                                                                       TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 18

CORRESPONDENCE ADDRESS:
ADDRESSEE: Mark M. Friedman c/o Anthony Casto
STREET: 2001 Jefferson Davis Highway, Suite 2
CITY: Arlington
STATE: Virginia
COUNTRY: United States of America
ZIP: 22202
COMPUTER READABLE FORM:
MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
COMPUTER: Twinhead* Slimnote-890TX
OPERATING SYSTEM: MS DOS version 6.2,
Windows version 3.11
                                                                                                                                                                                                         SOFTWARE: Word for Windows version 2.0 co to an ASCI file CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/383,630A FILING DATE: 26-Aug-1999 CLASSIFICATION: <Unknown> PRIOR APPLICATION DATA:
APPLICATION NUMBER: <Unknown> FILING DATE: <Unknown> FILING DATE: <Unknown> NAME: Friedmam, Mark M.
                                                                                                           NAME: Friedmam, Mark M.
REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 1402
TELECOMMUNICATION INFORMATION:
TELEPHONE: 972-3-562553
TELEFAX: 972-3-562554
TELEX: <Unknown >
  SEQUENCE
630-3
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                              STRANDEDNESS: single TOPOLOGY: linear
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                                                             amino
                                                           acid
                  SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Friedman c/o Anthony Castorina
son Davis Highway, Suite 207
                  ID
                  NO:
                                                                                                                                                                                 1402/2
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RESULT 10
US-09-173-151A-33
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; Sequence 33, Applica
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Best Local S
Matches 100
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GENERAL INFORMATION:
                                                                                                                   APPLICATION NUMBER: US 60/065,776
FILING DATE: 17-NOV-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/078,008
FILING DATE: 12-MAR-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/081,883
FILING DATE: 15-APR-1998
PRIOR APPLICATION NUMBER: US 60/095,987
FILING DATE: 10-AUG-1998
DBIOD ADDITION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 14-OCT-
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/173,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94304-1104
COMPUTER READABLE
MEDIUM TYPE: F
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               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 15-0CT-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER:
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Debets,
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Sana, Theodore R
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IBM PC compatible
NYSTEM: PC-DOS/MS
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California Avenue
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14-OCT-1998
     MBER: US 6
15-OCT-1997
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Best Local S
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TITLE OF INVE
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NUMBER OF SEC
CORRESPONDENC
ADDRESSEE:
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COMPUTER READAB
MEDIUM TYPE:
APPLICATION NU
FILING DATE:
CLASSIFICATION
ATTORNEY/AGENT I
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                                                                                                                                                                                                                                                    CITY:
STATE:
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REGISTRATION NUMBER: 34,090
REFERENCE/DOCKET NUMBER: DX07
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650)852-9196
TELEFAX: (650)496-1200
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 802 amino acids
                 CURRENT APPLICATION NUMBER: US/07/640 020 FILING DATE: 1991011
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INFORMATION:
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                                                                                                                                                                                                                               E: Chiron Corporation
4560 Horton Street
neryville
California
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larity 31.3%;
Conservative 5
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PE: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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Valenzuela, Pabio ....
Barr, Philip J.
Barr, Philip J.
FACTION: Expression and Use of Human Fibroblast
FENTION: Factor Receptor
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Valenzuela, Pablo D.T
INFORMATION:
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                                                            US/07/640,029
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; Pred. No. 2.2e-26;
50; Mismatches 136;
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Matches 89
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APPLICANT: IGARAS
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 816 amino acid
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MOLECULE TYPE:
7-640-029-1
FILING DATE: 16-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: LINEK, Ernest V
                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS SOFTWARE: PatentIn Release CURRENT APPLICATION DATA:
                                                                       APPLICATION NUMBER: 1
FILING DATE: 06-JUN-
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                     COMPUTER READABLE MEDIUM TYPE: F
                                                                                                                                                                                                                                                                            TITLE OF INVENTION: PR
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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CITY: BC
STATE: M
COUNTRY:
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                                   APPLICATION NUMBER: FILING DATE: APPLICATION NUMBER:
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US
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SENOO, Mas
                                                                                                                                                                                                                                                                                                                WATANABE,
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linear
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CUSHMAN
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BM PC compatible
PC-DOS/MS-DOS
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                                                                                   06-JUN-1995
V: 435
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Masaharu
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                                                           US/08/149,664
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RESULT 13
US-08-471-570-
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TELEPHONE: (617)523-3400
TELEFAX: (617)523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amino acids
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Best Local S
Matches 97
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                                                                                                                                                                                                                                                                                                                     Patent No.
                                                                                                                                                                                       APPLICANT: SENCE APPLICANT: WATANABE, APPLICANT: WATANABE, TITLE OF INVENTION: PROTINUMBER OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: DAVID G. COMMAN
                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: IGARAS
                                                                                                              ZIP: 02109
COMPUTER READAB
MEDIUM TYPE:
                                                          COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0,

CURRENT APPLICATION DATA:
PRIOR APPLICATION
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TOPOLOGY: lingle
                        APPLICATION FILING DATE: CLASSIFICATI
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CITY: Bost
STATE: Mass
                                                                                                                                                     COUNTRY:
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WATANABE, Tatsuya
ÆNTION: PROTEIN,
             TION: 435
ATION DATA:
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US
                                                                                                                                                                                           30 Water Street
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                                                  NUMBER:
  NUMBER:
                 06-JUN-1995
N: 435
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REGISTRATION NUMBER: 29822
REFERENCE/DOCKET NUMBER: 4089
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)523-3400
TELEFAX: (617)523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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Best Local
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GENERAL INFORMATION:
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08-471-570-
                                                                                                                                                  TITLE OF INVENTION: PROPERTY OF SEQUENCES: 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: DAVID G. CO
                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 0 FILING DATE: 16-AUG-1991 ATTORNEY/AGENT INFORMATION:
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                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0,

CURRENT APPLICATION DATA:
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APPLICATION FILING DATE:
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             NUMBER:
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Pred. No. 3.1e
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Best Local
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REGISTRATION NUMBER: 29822
REFERENCE/DOCKET NUMBER: 4089
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)523-3400
TELEFAX: (617)523-6440
TELEX: 200291 STRE UR
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: linear
MOLECULE TYPE: prote
)8-471-570-8
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
               ZIP: 19426
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                               NUMBER OF SEQUENCES: CORRESPONDENCE ADDRE
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INFORMATION:
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IBM PC compatib
SYSTEM: PC-DOS/
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larity 30.1%;
Conservative 59
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Crumley, Greg
Jaye, Michael C
Schlessinger, Joseph
ZENTION: Fibroblast Growth Factor
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16-AUG-1991
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; Pred. No. 4e-26;
59; Mismatches 129;
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Search completed: February Job time : 29 secs
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FILING DATE: 14-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/934,372
FILING DATE: 21-AUG-1992
PRIOR APPLICATION NUMBER: US 07/549,587
FILING DATE: 06-JUL-1990
ATTORNEY/AGENT INFORMATION:
NAME: Savitzky, Martin
REGISTRATION NUMBER: 29,699
REFERENCE/DOCKET NUMBER: A0496E
TELEPHONE: (610) 454-3816
TELEPHONE: (610) 454-3808
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 821 amino acids
TYPE: amino acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                            Query Match 21.9%;
Best Local Similarity 31.8%;
Matches 90; Conservative 5
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APPLICATION NUMBER: US
FILING DATE: 26-MAY-19
CLASSIFICATION: 435
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TOPOLOGY: linear
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26-MAY-1995
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Pred. No. 1e-25;
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RESULT 1
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fibroblast growth factor receptor ...
N; Contains: fibroblast growth factor receptor ...
C; Species: Xenopus laevis (African clawed frog)
C; Species: Xenopus laevis (African clawed frog)
C; Accession: A36477; C39752
R; Musci, T.J.; Amaya, E.; Kirschner, M.W.
Proc. Natl. Acad. Sci. U.S.A. 87, 8365-8369, 1990
A; Title: Regulation of the fibroblast growth factor receptor in (A; Reference number: A36477; MUID:91045998; PMID:2172985
A; Cross-references: UNIPROT:P22182; GB:U24491; GB:M37201; NID:91
A; Note: 45-Arg, 49-Thr, 50-Val, and 64-Ser also found
R; Friesel, R.; Dawid, I.B.
Mol. Cell. Biol. 11, 2481-2488, 1991
A; Title: cDNA cloning and developmental expression of fibroblas A; Arcession: C39752
A; Accession: C39752
A; A;Accession: C39752
A;Accession: C39752
A;Molecule type: mRNA
A;Residues: 1-30,119-189,'G',191-418,'L',420-636,'R',638-788,'V',790-812 <FRI
A;Residues: 1-30,119-1812,'Risk factor receptor 1; immunoglobulin homo
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglocation; glyc
protein kinase
F;12-17/Domain: signal sequence #status predicted <SIG>
F;22-372/Domain: extracellular #status predicted <EXT>
F;22-372/Domain: immunoglobulin homology <IM1>
F;125-132/Region: acidic
F;167-228/Domain: immunoglobulin homology <IM2>
F;333-393/Domain: transmembrane #status predicted <INM>
F;394-812/Domain: protein kinase homology <IM3>
F;394-812/Domain: protein kinase homology <IM3>
F;470-755/Domain: protein kinase homology <INN>
F;478-486/Region: protein kinase homology <KIN>
F;478-186/Region: protein kinase homology <KIN>
F;478-186/Region: protein kinase homology <KIN>
F;508,525,617/Active site: Lys, Glu, Asp #status predicted
F;622,635/Binding site: magnesium (Asn, Asp) #status predicted
F;622,635/Binding site: magnesium (Asn, Asp) #status predicted
F;648/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #stat Ş Query Ma Best Loc Matches Loca. sst growth factor receptor A2 precursor - African clawed frogins: fibroblast growth factor receptor A2, short splice form; ser Xenopus laevis (African clawed frog)
08-Mar-1991 #sequence_revision 19-Jan-1996 #text_change 09-Jusion: A36477; C39752 Match 106; 9 LLLPPLLLGAFPPAAAARGPPKMADKVVPR--awid, I.B. 11, 2481-2488, 1991 11, and developmental expression of fibroblast oning and developmental expression of fibroblast er: A39752; MUID:91203867; PMID:1850097 24.8%; Score 425.5; D larity 33.2%; Pred. No. 3.7e-Conservative 61; Mismatches B -23; 123; -QVARLGRTVRLQCPVEGDPPPLT 1; Length 812; Indels NID: 9857677; PIDN: AAA86868.1; ב short splice early 09-Jul-2004 29; growth (Asn) protein-tyrosine Gaps Xenopus factor #status homology; protes glycoprotein; gs #status <FRI> (covalent) 64 form #status embryos. receptors predicted predicte #8ta χir

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RESULT 2
A39752
Eibroblast growth factor receptor Al precursor - African clawed frog fibroblast growth factor receptor Al, short splice form; pro C; Species: Xenopus laevis (African clawed frog)
C; Date: 0.7-Feb-1992 #sequence_revision 19-Jan-1996 #text_change 09-Jul-: C; Date: 0.7-Feb-1992 #sequence revision 19-Jan-1996 #text_change 09-Jul-: 0.7-Feb-1992 #sequence revision 19-Jan-1996 #text_change 09-Jul-: 0.7-Feb-1992 #sequence revision 19-Jan-1996 #text_change 09-Jul-: 0.7-Feb-1992 #sequence revision 19-Jan-1996 #sequence receptor Al, short spire receptor Al, short spire receptor 19-Jan-1996 #sequence receptor Al, short spire receptor 19-Jan-1996 #sequence receptor 19-Jan-1996 #sequenc
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- TNRTRITGEEIQISNAGPEDNGLYACVTIGPSGTYTVLFS
                                 GWSRFRVLPQGLKVKQVEREDAGVYVÇKATNGFGSLSVNYTLVVLDDIS
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RESULT 3
B49151
fibroblast growth facto
C; Species: Pleurodeles
C; Date: 19-Dec-1993 #se
C; Date: 19-Dec-1993 #se
C; Accession: B49151
R; Shi, D.L.; Feige, J.J
Development 116, 261-27
A; Title: Differential e
A; Reference number: A49
A; Reference number: A49
A; Roccession: B49151
A; Status: preliminary
A; Molecule type: nuclei
A; Residues: 1-822 <SHI
A; Experimental source:
A; Note: Gemience extractions
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C; Superfamily: bas
C; Keywords: ATP; g
F; 283-354/Domain:
F; 484-769/Domain:
F; 492-500/Region:
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in: protein kinase homology <KIN>
n: protein kinase ATP-binding motif
VNGSRYGPDGVPFVQVLKTADINSSEVEVLYLHNVSFEDAGEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     e, J.J.; Riou, J.F.; DeSimone, D.W.; BOUCGUL, J.C.
261-273, 1992
tial expression and regulation of two distinct fibroblast
r: A49151; MUID:93130775; PMID:1483392
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deles waltlii (Iberian ribbed newt)
93 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
51
                                                            PPPLTMW-TKDGRTIHSGWSRFRVLPQGLKVKQVEREDAGVYVCKATNGFGSLSVNY 112
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                                 YGAEG----RHNSTIDVGGQKFVVLPTGDVWSRPDGSY 319
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protei

growth

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receptor

Iberian

ribbed newt

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C;Species: Pleurodeles waltlii (Iberian ri
C;Date: 06-Jan-1995 #sequence_revision 06-
C;Accession: S19947
R;Shi, D.L.; Feige, J.J.; Riou, J.F.; DeSi
submitted to the EMBL Data Library, March
A;Description: Receptores during early dev
A;Reference number: S19947
A;Reference number: S19947
A;Rocession: S19947
A;Rolecule type: mRNA
A;Residues: 1-822 <SHI>
A;Cross-references: UNIPROT:Q91288; EMBL:P
C;Superfamily: basic fibroblast growth fac
C;Keywords: ATP; growth factor receptor
F;283-354/Domain: immunoglobulin homology
F;484-769/Domain: protein kinase homology
TVHU2F
fibroblast growth factor receptor flg-2 pr
fibroblast growth factor receptor flg-2 pr
C; Species: Homo sapiens (man)
C; Date: 31-Dec-1993 #sequence_revision 31-
C; Accession: A60350; S21843
R; Avivi, A.; Zimmer, Y.; Yayon, A.; Yarden
Oncogene 6, 1089-1092, 1991
A; Title: Flg-2, a new member of the family
A; Reference number: A60350
A; Molecule type: mRNA
A; Residues: 1-800 <AVI>
A; Cross-references: EMBL:X58255; NID:g3138
A; Cross-references: EMBL:X58255; NID:g3138
A; Cross-references: GDB:T7273; OMIM:17694
A; Gene: GDB:FGFR2; JWS; CFD1; KGF; FLG2
A; Cross-references: GDB:127273; OMIM:17694
A; Map position: 10q25.3-10q26
C; Superfamily: basic fibroblast growth fac
C; Keywords: ATP; autophosphorylation; dupl
F;1-21/Domain: signal sequence #status pre
F;22-800/Product: fibroblast growth factor;
F;22-369/Domain: extracellular #status pre
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      Superfamily: basic fibroblast growth factor receptor 1; immuno (eywords: ATP; autophosphorylation; duplication; glycoprotein; (eywords: ATP; autophosphorylation; duplication; glycoprotein; -21/Domain: signal sequence #status predicted <SIG> .-21/Domain: fibroblast growth factor receptor flg-2 #statu?2-369/Domain: extracellular #status predicted <EXT>
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Library, March 1992
during early developm
                                                                                                                                                                                                                                            c of the family MUID:91296390;
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F;131-137/Region: a
F;262-335/Domain: i
F;370-390/Domain: t
F;391-800/Domain: i
F;464-749/Domain: i
F;472-480/Region: I
F;59-107,170-222,20
F;96,219,256,288,30
F;96,219,611/Activ
F;616,629/Binding site
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A; Note: sequence e
C; Superfamily: bas
C; Keywords: ATP; g
F; 262-335/Domain:
F; 464-749/Domain:
F; 472-480/Region:
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A48991
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A;Status: preliminary
A;Molecule type: nucl
                                                                                                                                                                                                                                                                                                                                                           C;Accession: A48991
R;Katoh, O.; Hattori, Y.; Sasaki, H.; Sakamoto, H.; Fujimoto,
Cancer Res. 53, 1136-1141, 1993
A;Title: Isolation of the complementary DNA encoding a mouse
A;Reference number: A48991; MUID:93177694; PMID:8382556
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C;Date: 19-Dec-19
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N;Contains:
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fus musculus (house mouse)
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GTGLVAS-HRILVGPQRLQVLNASHEDAGVYSCQHRLTR--
                                                        AGATSEPPGPEQRVVRRAAEVPGPEPSQQEQVAFGSGDTVELSCHPPGGAPTGPTVWAKD
                                                                                          AAAARGPPKMADKVVPR----
                                                                                                                                                                                                                                                                      source: brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    growth factor receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23.8%; ilarity 34.3%; Conservative 48
                                                                                                                                                                           extracted from NCBI backbone (NCBIN:126536, NCBIP:126537)
asic fibroblast growth factor receptor 1; immunoglobulin homo-
growth factor receptor; heparin binding; phosphotransferase;
immunoglobulin homology <IMM>
protein kinase homology <KIN>
protein kinase ATP-binding motif
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protein kinase ATP-binding motif
269-333/Disulfide bonds: #status predicted
309,322/Binding site: carbohydrate (Asn) (covalent) #st
ive site: Lys, Glu, Asp #status predicted
site: magnesium (Asn, Asp) #status predicted
e: phosphate (Tyr) (covalent) (by autophosphorylation)
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                                                                                                                                                                                                                                                                                      UNIPROT: Q61851; GB: S56291; NID: g298329;
                                                                                                                                                                                                                                                                                                                                                                                                                                       #sequence
                                                                                                                                                                                                                                                                                                                     acid
                                                                                                                                    23.8%;
                                                                                                                                                                                                                                                                                                                                                                                                         Sagaki, H.; Sakamoto, H.; Fujimoto,
                                                                                                                                                                                                                                                                                                                                                                                                                                        revision 18-Nov-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 409; DB 1; Le
Pred. No. 5.8e-22;
48; Mismatches 111;
                                                                                                                     Score 408.5; DE
Pred. No. 6.3e-2
19; Mismatches 1
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                                                                                        -QVA-RLGRTVRLQC-PVEGDPPPLTMWTKD
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114;
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    -RVLCHFSVRVTDAPSSGD
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RESULT 7

155363

fibroblast growth factor receptor 3 - mouse C;Species: Mus musculus (house mouse)
C;Date: 02-Jul-1996 #sequence_revision 02-Jul-1996 C;Accession: I55363; BS3627
R;Ornitz, D.M.; Leder, P.
J. Biol. Chem. 267, 16305-16311, 1992
A;Title: Ligand specificity and heparin dependence A;Reference number: I55363; MUID:9235591; PMID:137
A;Accession: I55363
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-801 <RES>
A;Cross-references: UNIPROT:Q61851; GB:M81342; NID:A;Cross-references: UNIPROT:Q61851; GB:M81342; NID:A;Cross-references: UNIPROT:Q61851; GB:M81342; NID:A;Cross-references: LMIPROT:Q61851; GB:M81342; NID:A;Cross-references: A53627; MUID:94209351; PMID:751
A;Accession: B53627
A;Accession: B53627
A;Accession: B53627
A;Accession: B53627
A;Accession: B53627
A;Accession: B53627
A;Accession: B33627
A;Cross-references: GB:L26492
C;Genetics:
A;Gene: mFR3
A;Introns: 304/3; 353/1
C;Superfamily: basic fibroblast growth factor receptor F;262-335/Domain: immunoglobulin homology <NIM>F;262-335/Domain: protein kinase ATP-binding motif F;472-480/Region: protein kinase ATP-binding motif
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$38579
fibroblast growth
C; Species: Pleurod
C; Date: 06-Jan-1991
C; Accession: $38579
R; Shi, D.L.; Fromes
submitted to the Electric State S
                                                                            A;Status: prelim
A;Molecule type:
A;Residues: 1-80
A;Cross-referenc
C;Genetics:
A;Gene: cek2
C;Superfamily: b
C;Keywords: ATP;
F;1-23/Domain: s
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A; Cross-references: U
C; Superfamily: basic
C; Keywords: ATP; grov
F; 258-331/Domain: imn
F; 456-741/Domain: pro
F; 464-472/Region: pro
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C;Species: Gallus g
C;Date: 09-Nov-1990
C;Accession: A35963
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A; Reference number:
A; Accession: S38579
A; Status: prelimina
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A35963
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R;Pasquale, E.B.
Proc. Natl. Acad. Sci. U.S.A. 87,
A;Title: A distinctive family of
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s: UNIPROT:P18460;
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gallus (chicken)
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GB:M35195;

NID: 9211442;

PIDN: AAA48664.1;

PID:g211443

immunoglobulin homology; protei
otein; growth factor receptor;

protein-tyrosine PMID:2165604

kinase

receptors

#text_change

09-Jul-2004

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basic fibroblast growth factor receptor
p; growth factor receptor
in: immunoglobulin homology <IMM>
in: protein kinase homology <KIN>
on: protein kinase ATP-binding motif
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Expression of FGFR-3 in amphibian embryos
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TYMUR3
fibroblast growth factor receptor 3 precursor - human
N; Contains: protein-tyrosine kinase (EC 2.7.1.112)
C; Species: Homo sapiens (man)
C; Date: 31-Dec-1993 #sequence revision 31-Dec-1993 #text_change
C; Accession: A38576; A55273; E38289; IS1880
R; Keegan, K.; Johnson, D.E.; Williams, L.T.; Hayman, M.J.
Proc. Natl. Acad. Sci. U.S.A. 88, 1095-099, 1991
A; Title: Isolation of an additional member of the fibroblast gro
A; Reference number: A38576; MUID:91142118; PMID:1847508
A; Accession: A38576
A; MOID:91142118; PMID:1847508
A; Accession: A38576
A; MOID:91142118; PMID:1847508
A; Residues: 1-806 <KEE>
A; Thompson, L.M.; Plummer, S.; Schalling, M.; Altherr, M.R.; Gus
Genomics 11, 1133-1142, 1991
A; Title: A gene encoding a fibroblast growth factor receptor iso
A; Reference number: A55273; MUID:92147110; PMID:1644411
A; Accession: A55273
A; Status: nucleic acid sequence not shown
A; Residues: nucleic acid sequence not shown
A; Residues: nucleic acid sequence not shown
A; Residues: 1-6-394; V', 396-806 <THO
A; Cross-references: GB:M64347, NID:9182564; PIDN:AAA58470.1; PID
A; Rossiues: 10; Mackelae, T.P.; Alitalo, R.; Lehvaeslaiho, H.; A
Proc. Natl. Acad. Sci. U.S.A. 87, 8913-8917, 1990
A; Title: Putative tyrosine kinases expressed in K-562 human leuk
A; Reference number: A38268; MUID:91062389; PMID:2247464
A; Accession: E38269
A; MOID:91062389; PMID:2247464
A; Accession: E38269
A; MOID:91062389; PMID:2247464
A; Accession: GB:M37782
R; Bellus, G.A.; Hefferon, T.W.; Ortiz de Luna, R.I.; Hecht, J.T.
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F;3131-138/Region: acidic F;262-335/Domain: immunoglobulin homology <IMM>
F;369-389/Domain: transmembrane #status predicted <TMM>
F;369-389/Domain: intracellular #status predicted <INT>
F;464-749/Domain: protein kinase homology <KIN>
F;472-480/Region: protein kinase homology <KIN>
F;61-107,170-222,269-333/Disulfide bonds: #status predicted F;61-219,256,288,309,322/Binding site: carbohydrate (Asn) (cova F;616,629/Binding site: magnesium (Asn, Asp #status predicted F;616,629/Binding site: phosphate (Tyr) (covalent) (by autophosphone)
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i0, 1989 ation and complementary DNA Ar: A41345; MUID:89298406; I

cloning of a PMID:2544996

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A; Map position: 4p16.3-4p16.3

C; Function:
A; Description: receptor for both acidic and basic fibroblast growth factors
A; Description: receptor for both acidic and basic fibroblast growth factor receptor 1; immunoglobulin homolog
C; Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homolog
C; Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homolog
C; Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homolog
C; Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin prowth factor receptor 3 #status predicted <SIG>
F; 1-22/Domain: signal sequence #status predicted <SIG>
F; 23-375/Domain: extracellular #status predicted <EXT>
F; 33-319/Region: acidic
F; 36-341/Domain: immunoglobulin homology <IMM>
F; 376-396/Domain: intracellular #status predicted <INT>
F; 470-755/Domain: protein kinase homology <KIN>
F; 478-486/Region: protein kinase homology <KIN>
F; 478-486/Region: protein kinase ATP-binding motif
F; 61-109, 176-228, 275-339/Disulfide bonds: #status predicted
F; 98, 225, 262, 294, 315, 328/Binding site: carbohydrate (Asn) (covalent) #status predicted
F; 622, 635/Binding site: magnesium (Asn, Asp) #status predicted
F; 648/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status
Accession: A41345; MUID: 892984(A;Residue: mucleic acid sequent
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A; Cross-references
A; Map position: 4p
C; Function:
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A;Molecule type:
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A;Note: this sequ
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cences: GB:S76733; NID:g914201; PIDN:AAB33323.1; PID:g914202
sequence represents a mutant form associated with achondroplasia
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plasia is defined by recus
r: 151880; MUID:95150025;
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larity 33.4%; Pred. No. 8.4e-21;
Conservative 49; Mismatches 116
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Proc. Natl. Acad. Sci. U.S.A. 86, 5449-5453, 1989

A;Title: Identification of a developmentally regulated protein-tyrosine kinas A;Reference number: A33908; MUID:89315814; PMID:2473471

A;Residues: A33908; MUID:89315814; PMID:2473471

A;Residues: 1-89,'A',91-685,'M',687-819 <PAS>
A;Cross-references: GB:M24637

A;Note: this protein is expressed in embryonic tissues and, at low levels, ir c;Genetics:
A;Gene: cekl
C;Genetics:
A;Gene: cekl
C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin home C;Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor F;1-21/Domain: signal sequence #status predicted <SIG>F;2-319/Product: fibroblast growth factor receptor 1 #status predicted <MAT: F;2-374/Domain: extracellular #status predicted <EXT>
F;22-374/Domain: extracellular #status predicted <TMM>F;395-395/Domain: immunoglobulin homology <IMM>F;395-395/Domain: intracellular #status predicted <INT>F;474-759/Domain: protein kinase homology <KIN>F;482-490/Region: protein kinase homology <KIN>F;482-490/Region: protein kinase ATP-binding motif F;5116,225,238,262,294,315,339/Binding site: carbohydrate (Asn) (covalent) F;512,529,621/Active site: Lys, Glu, Asp #status predicted F;626,639/Binding site: magnesium (Asn, Asp) #status predicted from the protein kinase homology is the carbohydrate (Asn) (covalent) F;652/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #status predicted
RESULT 12

B54846
fibroblast growth factor receptor b precursor - C; Species: Rattus norvegicus (Norway rat)
C; Date: 21-Jul-1995 #sequence_revision 28-Jul-1
C; Accession: B54846
R; Takagi, Y.; Shrivastav, S.; Miki, T.; Sakaguc
J. Biol. Chem. 269, 23743-23749, 1994
A; Title: Molecular cloning and expression of th
ility and covalent attachment of heparan sulfat
A; Reference number: A54846; MUID: 94375484; PMID
A; Accession: B54846
A; Status: preliminary; not compared with concep
A; Molecule type: mRNA
A; Residues: 1-822 < TAK>
C; Superfamily: basic fibroblast growth factor r
C; Keywords: ATP; growth factor receptor
F; 172-233/Domain: immunoglobulin homology < IMM;
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A;Note: part
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A;Introns: 43/1; 125/1; 158/1; 213/1; 254/1; 318/1; 364/1; 429/1; 478/1; 556/1; 619/1; C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prot F;1-26/Domain: signal sequence #status predicted <SIG>
F;27-818/Product: fibroblast growth factor receptor-4 #status predicted <MAT>
F;27-818/Product: fibroblast growth factor receptor-4 #status predicted <MAT>
F;37-238/Domain: immunoglobulin homology <IM1>
F;177-238/Domain: immunoglobulin homology <IM2>
F;276-347/Domain: immunoglobulin homology <IM2>
F;382-402/Domain: transmembrane #^-
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F;485-493/Po--
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C; Date: 29-Jun-1995
C; Accession: JC4058
R; Shiozaki, C.
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lloning of cDNA and genomic DNA encoding fibroblast growth e number: JC4058; MUID:95137391; PMID:7835703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ASTVVGGDVEFVCKVYSDAQPHIQWIKHVEKNGSKYGPDGLPYLKVLKHSG---INSSNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PPTKYQISQPE-ACVVAP
                                                                                                                                                    22.5%; Score 385.5; DB larity 29.5%; Pred. No. 3e-20; Conservative 55; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     VRNQHWSLIMESVVPSDKGNYTCLVENEYGSINHTYHLDVVERSPHRPILQAGLPAN
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                                                            TIHSGWSRFRVLPQG--
                                                                                                                        3AFPPAAAARGPPKMAD--
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: UNIPROT:Q91742; DDBJ:D31761; NID:g809527; PIDN:BAA06539.1; PID:g80952
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   actor receptor-4 precursor - African clawed frog
laevis (African clawed frog)
#sequence_revision 14-Jul-1995 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22.7%;
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kinase ATP-binding motif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 389.5; DB 2;
Pred. No. 1.5e-20;
56; Mismatches 132;
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                                                                                                                       ----KVVPRQVARLGRTVRLQCPVEGDPPPLTMWTK
                                                     LKVKQVEREDAGVYVCKATNGFGSLSVNYTLVVL
  -QWARPRFTQPSKMRRRVIARPVGSSVRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Saigo, K.; Emori, Y.; Shiokawa,
                                                                                                                                                                                   DB 2;
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H.

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RESULT 14

$24108

$24108

protein-tyrosine kinase (EC 2.7.1.112) bek - chicken

protein-tyrosine kinase (EC 2.7.1.112) bek - chicken

protein-tyrosine kinase (EC 2.7.1.112) bek - chicken

protein-tyrosine kinase bek

C;Specias: Gallus gallus (chicken)

C;Date: 25-Feb-1994 #sequence_revision 10-Nov-1995 #text_change 09-Jul-2004

C;Accession: S24108

R;Sato, M.; Kitazawa, T.; Iwai, T.; Seki, J.; Sakato, N.; Kato, J.; Takeya, T.

Oncogene 6, 1279-1283, 1991

A;Title: Isolation of chicken-bek and a related gene; identification of structural vari

A;Reference number: S24108

A;Reference number: S24108

A;Reference number: S24108

A;Reference number: S24108

A;Reference protein from PMID: 91319411; PMID: 1650446

A;Residues: preliminary

A;Residues: 1-824 <SAT>

A;Cross-reference: UNIPROT: Q90749; EMBL: X61992; NID: g63085; PIDN: CAA43965.1; PID: g6308

C;Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin homology; prot

C;Keywords: ATP; growth factor receptor; phosphotransferase; tyrosine-specific protein

F;174-235/Domain: protein kinase homology <KIN>

F;482-767/Domain: protein kinase homology <KIN>

F;490-498/Region: protein kinase ATP-binding motif
RESULT 15
TVHUF4
fibroblast g
N;Alternate
N;Contains:
C;Species: H
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 growth factor receptor 4 names: protein-tyrosine protein-tyrosine kinase Homo sapiens (man)
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Similarity 30.0%;
%1; Conservative
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Pred.
                    precursor
kinase tkf
(EC 2.7.1.
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4.2e-20;
ches 114;
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                       112)
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---ALLGVLLSVPGPPVLSLEASEEVELEPCLAPSLEQQEQELTVALGQPVRLCC

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A; Description: receptor mediating effects of fibroblast growth factor
A; Note: expressed in normal lung; expressed in some carcinomas
C; Superfamily: basic fibroblast growth factor receptor 1; immunoglobulin home
C; Keywords: ATP; autophosphorylation; duplication; glycoprotein; growth factor
F; 1-24/Domain: signal sequence #status predicted <SIG>
F; 25-802/Product: fibroblast growth factor receptor 4 #status predicted <MAT:
F; 25-369/Domain: extracellular #status predicted <EXT>
F; 26-103/Domain: immunoglobulin homology <IM1>
F; 165-226/Domain: immunoglobulin homology <IM2>
F; 264-335/Domain: immunoglobulin homology <IM3>
F; 370-390/Domain: transmembrane #status predicted <INT>
F; 465-750/Domain: intracellular #status predicted <INT>
F; 465-750/Domain: protein kinase homology <KIN>
F; 473-481/Region: protein kinase homology <KIN>
F; 473-481/Region: protein kinase homology <KIN>
F; 473-481/Region: protein kinase homology <KIN>
F; 503,520,612/Active site: Lys, Glu, Asp #status predicted
F; 503,520,612/Active site: Lys, Glu, Asp #status predicted
F; 617,630/Binding site: magnesium (Asn, Asp) #status predicted
F; 643/Binding site: phosphate (Tyr) (covalent) (by autophosphorylation) #stat
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J. Biol. Chem.
A;Title: Fibroblast
A;Reference number:
A;Accession: A46615
A;Accession: nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 31-LEC
C;Accession: S1534!
R;Partanen, J.; Mar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Title: Two additional A; Reference number: S190 A; Recession: A41598 A; Status: nucleic acid s A; Molecule type: mRNA A; Residues: 399-534, 'M', A; Experimental source: R; Partanen, J.; Maekelas Proc. Natl. Acad. Sci. U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: 1
A: Residues: 1-802
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A;Gene: GDB
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: Fibroblast growth factor receptor 4 is a high affinity receptor for boence number: A46615; MUID:93194827; PMID:7680645
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1. Sci. U.S.A. 87, 8913-8917, 1990
ve tyrosine kinases expressed in K-562 human leukemia cells.
per: A38268; MUID:91062389; PMID:2247464
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X MEDLINE=22388257; PubMed=12477932;

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X MEDLINE=223882;

X MEDLINE=223882;

X MEDLINE=223882;

X MEDLINE=2238
                                                           "The secreted protein discovery i effort to identify novel human se bioinformatics assessment."; Genome Res. 13:2265-2270(2003).

EMBL; BC036769; AAH36769.1; -.

EMBL; AY358303; AAQ88670.1; -.

HSSP; P11362; 1EVT.

Genew; HGNC:3693; FGFRL1.

GO; GO:0004872; F:receptor activi
InterPro; IPR007110; Ig-like.

Pfam; PF00047; ig; 3.

PROSITE; PS50835; IG_LIKE; 3.
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                                                                                                                                                                                                                                                                                                                                                 Godowski P.;
"The secreted effort to ide
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SEQUENCE FROM TISSUE=Brain;
Strausberg R.;
Submitted (AUG
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Huang A.,
Lewis L.,
Seshagiri
Vandlen R.
Yi S., Yu
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Q8N441;
01-OCT-2002
01-OCT-2002
01-OCT-2004
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SEQUENCE
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01-OCT-2004 (TrEMBLrel. 28, Last annot)
Fibroblast growth factor receptor-like
Name=FGFRL1; ORFNames=UNQ480;
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Proc. Natl.
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, Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J., S., Simmons L., Singh J., Smith V., Stinson J., Vagts A., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D., G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA sequences."
L. Acad. Sci. U.S
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Best Loo
Matches
                                                                                                MEDLINE=21167383; PubMed=11267671;
Kim I., Moon S.O., Yu K.H., Kim U.H., Koh G.Y.;
"A novel fibroblast growth factor receptor-5 preferenti
in the pancreas.";
Biochim. Biophys. Acta 1518:152-156(2001).
EMBL; AJ277437; CAC14171.1; -.
EMBL; AF279689; AAK26742.1; -.
HSSP; P11362; 1EVT.
GO; GO:0016021; C:integral to membrane; NAS.
GO; GO:0005007; F:fibroblast growth factor receptor act
GO; GO:0001558; P:regulation of cell growth; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003598; Ig_c2.
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Mammalia; Eut
NCBI_TaxT
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Q9H4D7;
01-MAR-2001
01-MAR-2001
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FGFR-like pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM
TISSUE=Cartil
PubMed=110311
Receptor;
SIGNAL
SEQUENCE
                                                       SMART; SN PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                         Genomics [2]
                                                                                                                                                                                                                                                                                                                                                                                                          Wiedemann M., Trueb B.;
"Characterization of a novel protein (FGFRL1) from human cartilage related to FGF receptors.";
Genomics 69:275-279(2000).
                                                                                             InterPro; IPR
Pfam; PF00047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=FGFRL1
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(Fibroblast growth factor
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0; Mismatches 1;
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K PubMed=12975309;

K Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,

A Clark H.F., Gurney A.L., Abaya E., Baker K., Baldwin D., Brush J.,

A Clark H.F., Gurney A.L., Crowley C., Currell B., Deuel B., Dowd P.

A Chen J., Chow B., Chui C., Crowley C., Currell B., Deuel B., Dowd P.

A Chen J., Foster J., Grimaldi C., Gu Q., Hass P.E., Heldens S.,

A Huang A., Kim H.S., Klimowski L., Jin Y., Johnson S., Lee J.,

A Lewis L., Liao D., Mark M., Robbie E., Sanchez C., Schoenfeld J.,

Seshagiri S., Simmons L., Singh J., Smith V., Stinson J., Vagts A.,

A Vandlen R., Watanabe C., Wieand D., Woods K., Xie M.H., Yansura D.,

A Yi S., Yu G., Yuan J., Zhang M., Zhang Z., Goddard A., Wood W.I.,

Godowski P.;

T "The Secreted Protein Discovery Initiative (SPDI), a Large-Scale

Effort to Identify Novel Human Secreted and Transmembrane Proteins:

Bioinformatics Assessment.";

Bioinformatics Assessment.";
                                                                                                                 Query Ma
Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens (Human)
Eukaryota; Metazoa; (
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ88670;
AAQ88670;
02-MAR-2004
02-MAR-2004
02-MAR-2004
FGFRL1:
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hes 323; Conservative
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2e-124;
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Best Local
Matches 2
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Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; C Mammalia; Eutheria; Rodentia; S NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Q7TQM3
Q7TQM3;
01-OCT-2003
01-OCT-2003
01-MAR-2004
                                                                                                                                                                                                                                                                                                                                                               Receptor;
SIGNAL
CHAIN
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SEQUENCE FROM N.A.
STRAIN=Sprague-Dawley;
PubMed=12813049;
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Signal.
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growth factor rec
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529
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J. Biol. Chem. 278:33857-33865(2003).

EMBL; AJ536020; CAD59914.1; -.

GO; GO:0004872; F:receptor activity; IEI
InterPro; IPR007110; Ig-like.
InterPro; IPR003598; Ig_c2.

Pfam; PF00047; ig; 3.

SMART; SM00408; IGc2; 3.

PROSITE; PS50835.
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                                                                                                SPLLLLLPPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPL
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25, Last sequence update)
26, Last annotation update)
receptor-like protein prec
                                                                                                                                                                               Score 1561; DB 2;
Pred. No. 2e-112;
); Mismatches 15;
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fibroblast growth factor
protein.
page 23D67B419335FFFC CRC64;
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CTISSUE=Mammary tumor. ru...
CDiopsy:
CDiopsy:
CDiopsy:
XMEDLINE=22388257; PubMed=12477932;
XMEDLINE=22388257; PubMed=12.A., Grouse L.H., Schaefer C.F., Bhat N.K.
Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.
Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,
RA Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz
RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange
RA Brownstein M.J., WcKernan K.J., Abramson R.D., Mullah,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne |
RA Bosak S.A., MocEwan P.J., McKernan K.J., Malek J.A., Gibbs R.A.,
Mingny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Mingny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
Mingny D.M., Sodergren E.J., Rodrigues S., Sanc
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STRAIN=BALB/cJ;
MEDLINE=21311632; PubMed=1
Sleeman M., Fraser J., McD
Kumble K., Watson J.D., McD
Kumble K., Watson of a new f
"Identification of a new f
Gene 271:171-182(2001).
[2]
SEQUENCE FROM N.A.
STRAIN=BALB/cJ;
Sleeman M.A., Fraser J.K.,
Submitted (NOV-2000) to th
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STRAIN=BALB /c; TISSU
Wiedemann M., Trueb B
Submitted (DEC-2002)
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"Characterization of
related to FGF reco
Genomics 69:275-279
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SEQUENCE FROM N.
STRAIN=FVB/N-3;
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Q91V87;
O1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Fibroblast growth factor receptor 5 beta precursor (FGF protein 1 precursor) (Fibroblast growth factor receptor-Name=Fgfrl1; Synonyms=FGFRL1, Fgfr5;
Mus musculus (Mouse).
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ALB /c; TISSUE=Liver;
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5-279(2000)
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er J., McDonald M., Y
n J.D., Murison J.G.;
of a new fibroblast g
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EMBL/GenBank/DDBJ (
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M., Schuler G.,
C.F., Bhat N.K.
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RESULT 7
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AC AAH58
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COS Mus m
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(Tremblrel. 27, Created)
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rowth factor receptor-like 1.

PRELIMINARY;

529

AAH58745 PAAH58745;
02-MAR-2004 ('02-MAR-2004 ('02-MAR-2004 ('Fibroblast grown musculus Eukaryota; Memmalia; Eutincell_TaxID=10

letazoa; theria;

Chordata; Rodentia;

Craniata; Vertebrata; i Sciurognathi; Muridae;

Euteleostomi; Murinae; Mus

(Mouse)

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Strausberg R.;
Strausberg R.;
Submitted (SEP-2003) to the EM
EMBL; AF321300; AAL06295.1; -.
EMBL; AJ293947; CAC82376.1; -.
EMBL; BC058745; AAH58745.1; -.
EMBL; BC058745; AAH58745.1; -.
HSSP; P11362; 1EVT.
MGD; MGI:2150920; Fgfrll.
GO; GO:0005886; C:plasma membra
GO; GO:0005515; F:protein bindi
GO; GO:0008285; P:negative regu
InterPro; IPR007110; Ig-like.
Pfam; PF00047; ig; 3.
PROSITE; PS50835; IG_LIKE; 3.
Receptor; Signal.
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SIGNAL
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SEQUENCE
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886; C:plasma membrane; IDA.
515; F:protein binding; IPI.
5. B. negative regulation of c
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29 AA;
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larity 91.4%;
Conservative
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Acad. Sci. U.S.A.
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I., Touchman J.W., Green E.D., Dickson M.C.,
I., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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57013 MW;
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                                                                                                                                                                                                                                                   Potential.
fibroblast growth factor receptor 6CE76A75BF2498DD CRC64;
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SEQUENCE FROM N.A.

STRAIN=FVB/N-3; TISSUE=Mammary tumor;

K STRAIN=FVB/N-3; TISSUE=Mammary tumor;

K MEDLINE=22388257; PubMed=12477932;

A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Liatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

A Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Barownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Wsdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Halkesley R.W., Touchman J.W., Green E.J., Dickson M.C.,

A Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

A Jones S. J. Marra M.A.:
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Matches
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"Generation and initial analysis of and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:10
[2]
SEQUENCE FROM N.A.
STRAIN=FVB/N-3; TISSUE=Mammary tume
Strausberg R.;
Submitted (SEP-2003) to the EMBL/GEMBL; BC058745; AAH58745.1; -.
Receptor.
SEQUENCE 529 AA; 57013 MW; 6CE
Q6ZMD4;
Q6ZMD4;
Q6ZMD4;
Q5-JUL-2004 (TrEMBLrel. 27, Cre
05-JUL-2004 (TrEMBLrel. 27, Las
05-JUL-2004 (TrEMBLrel. 27, Las
Hypothetical protein FLJ23990.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Primates; C
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Catarrhini;
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AC BAD2
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OS Homo
OC Euka
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REMBL; AK172829; BAD18794.1; -.

RGO; GO:0004872; F:receptor activity; IEA.

InterPro; IPR003599; Ig.

R InterPro; IPR003598; Ig-like.

R InterPro; IPR003598; Ig-like.

R InterPro; IPR003598; Ig-like.

R SMART; SM00409; IG; 3.

R SMART; SM00409; IG; 3.

R PROSITE; PS50835; IG_LIKE; 3.

R PROSITE; PS50835; IG_LIKE; 3.

Receptor.

3 SEQUENCE 440 AA; 47587 MW. PROSITE
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Eukaryota; Me
Mammalia; Eutl
NCBI_TaxID=96(
[1]
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                                                                                                                                                      FROM N.A.
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ilarity 99.7%;
Conservative (
                   90.4%;
ilarity 99.7%;
Conservative
                                                                                                                                                                                     Metazoa;
utheria;
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                                                                                                                                                                                                                   (TrEMBLrel. 27, Created)
(TrEMBLrel. 27, Last sequence update)
(TrEMBLrel. 27, Last annotation update)
(TrEMBLrel. 27, Last annotation update)
(Ofis, clone HRC08053, highly similar to Homo sapiens)
(FGFRL1).
                                                                                                                                                                                                                                                                                                                                          606;
                                                                                                                                                                           606;
                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                 40 AA;
                                                                                     29; BAD18794.1;
                                                                                                                                                                                                            (Human).
                                                                                                                                                                                     Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                 47587 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1553; DB 2;
Pred. No. 6.6e-112;
0; Mismatches 1;
                      0;
                      Score 1553; DB 2
Pred. No. 6.6e-11
); Mismatches
                                                                                                                                                                                                                                                                                      PRT;
                                                                  2B6E0C2285AC3B59 CRC64;
                                                                                                                                                                                                                                                                                      440 AA
                      DB 2;
6e-112;
s 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                             Length 440;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                         Indels
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Shibahara T.
                                                                                                                                   a Y., Okamo
Shibahara
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q920C2;

PRELIMINARY;

PRT;

Q920C2;

I 01-DEC-2001 (TrEMBLrel. 19, Created)

I 01-DEC-2001 (TrEMBLrel. 19, Last sequence update)

I 01-DEC-2001 (TrEMBLrel. 26, Last annotation update)

I 01-MAR-2004 (TrEMBLrel. 26, Last annotation update)

Fibroblast growth factor receptor 5 beta/gamma (Fragment).

Name=Fgfrll; Synonyms=Fgfr5;

Mus musculus (Mouse).

C Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleos

C Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae

X NCBI_TaxID=10090;

N [1]

P SEQUENCE FROM N.A.

C STRAIN=BALB/cJ;

X MEDLINE=21311632; PubMed=11418238;

I Identification of a new fibroblast growth factor receptor,

Gene 271:171-182(2001).
                                                                           Query Ma
Best Loo
Matches
                                                                                                                                                                                                                                                                   Gene 271:171-182(2001).

[2]

SEQUENCE FROM N.A.

STRAIN=BALB/cJ;

Sleeman M.A.; Fraser J.K., McDonald M.,

Grandison P., Kumble K.D., Watson J.D.,

Submitted (NOV-2000) to the EMBL/GenBank

EMBL; AF321302; AAL06297.1; -.

HSSP; P11362; 1EVT.

MGD; MGI:2150920; Fgfrll.

GO; GO:0005886; C:plasma membrane; IDA.

GO; GO:0005515; F:protein binding; IPI.

GO; GO:0008285; P:negative regulation of

InterPro; IPR003598; Ig-like.

InterPro; IPR003598; Ig-like.

SMART; SM00408; IGC2; 3.

PROSITE; PS50835; IG_LIKE; 3.
                                                                                                                                                                                                                                      Receptor
NON_TER
                                                                                                                                                            SEQUENCE
                                                                            Local S:
hes 281
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93.7%;
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Pred.
9; Mis
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fibroblast
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392A7D95
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growth
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.9e-108;
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Murinae; Mus
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NCB1-[1]

SEQUENCE FROM N.A.
TISSUE=Cartilage;
PubMed=12813049;
Trueb B., Zhuang L., Taeschler S., ...
"Characterization of FGFRL1, a Novel FG!
"Characterization of FGFRL1, a Novel FG!
T expressed in skeletal tissues.";
J. Biol. Chem. 278:33857-33865(2003).
DR GO; GO:0004872; F:receptor activity; I
DR InterPro; IPR007110; Ig-like.
DR InterPro; IPR003598; Ig_c2.
DR Pfam; PF00047; ig; 3.
SMART; SM00408; IGc2; 3.
PROSITE; PS50835; IG_LIKE; 3.
PROSITE; Signal.

1 18 fibrob]
Proteir
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Q7T2H2
ID Q7T2H2
AC Q7T2H2;
DT 01-OCT-20
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Best Local
Matches 2:
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Eukaryota; Met
Archosauria; A
Gallus.
NCBI_TaxID=903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-OCT-2003 (
01-MAR-2004 (
Fibroblast gr
Name=FGFRL1;
                                                                                                                                                                                                                                                                                                                                                                                                       Match
                                        183
                                                                               189
                                                                                                                       123
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238; C
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                                                                      QALTRPEAAEPRKKKWTLSLKNLRPEDSGKYTCRVSNRAGAINATYKVDVIQRTRSKPVL
                                                                                                                   -SEQ
                                                                                                                                                                                                                                                                                 LQLALLLAGIVALSDSARGPPRIADKVIHRQSVRLGRTIKLLCPVEGDPPPLTMWMKDGR
                                                                                                                                                                                                                                                                                                                           TTT
                                        KPLMPHEIGENKKKKWTLNLKNLKPEDSGKYTCRVFNKVGEINATYKVEVIQRTRSKPIL
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                                                                                                                                                                                                                                                                                                         PPLLLGAFPPAAAARGPPKMADKVVPRQVARLGRTVRLQCPVEGDPPPLTMWTKDGR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          etazoa;
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larity 75.8%;
Conservative 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Chicken).
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26, Last annotation update)
receptor-like protein precursor
                                                                                                                                                                                                                                                                                                                                                               Score 1306; DB 2;
Pred. No. 9.9e-93;
36; Mismatches 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                              Potential.
fibroblast growth factor receptor-
protein.
FFD0132AD917FF94 CRC64;
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Novel FGF recept
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Phasianinae;
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.

A Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunaratne P.H.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Nillalon D.K., Touchman J.W., Green E.D., Dickson M.C.,

Butterfield Y.
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"Generation and initial analysis of mor and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-[3]
Query
                                                                        EMBL; AJ574916; CAE00573.1; -.
EMBL; BC053245; AAH53245.1; -.
ZFIN; ZDB-GENE-040128-2; fgfrll.
GO; GO:0004872; F:receptor activity; I.
InterPro; IPR007110; Ig-like.
InterPro; IPR003598; Ig_c2.
Pfam; PF00047; ig; 3.
SMART; SM00408; IGc2; 3.
PROSITE; PS50835; IG LIKE.
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Actinopterygii; Neopterygii; Te
Cyprinidae; Danio.
NCBI_TaxID=7955;
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01-OCT-2003 (TrEMBLrel. 25, L
01-OCT-2004 (TrEMBLrel. 28, L
FGF receptor-like protein pre
receptor-like 1).
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Q7SX76;
01-OCT-2003
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FGF receptor
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TISSUE=Kidney;
Strausberg R.;
Submitted (JUN-2003) t
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J. Biol. C
[2]
SEQUENCE F
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                          Receptor;
SIGNAL
CHAIN
SEQUENCE
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MEDLINE=22388257;
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or; Signal.
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483
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  69.5%;
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                          Potential.
FGF receptor-like protein
3C2B34E5D57C332C CRC64;
  Score
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D.E., Schnerch A.,
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Ostariophysi; Cypriniformes;
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Matches
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Matches 211
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SEQUENCE FROM N.A.
STRAIN=BALB/cJ;
MEDLINE=21311632; PubMed=11418238;
Sleeman M., Fraser J., McDonald M., Yuar
Sleeman M., Watson J.D., Murison J.G.;
"Identification of a new fibroblast grounder 271:171-182(2001).
                                                                                                                                                                                                                                                                                                                                       Q920C3;
Q920C3;
Q920C3;
Q1-DEC-2001 (
Q1-DEC-2001 (
Q1-MAR-2004 (
Fibroblast gr
Name=Fgfrl1;
                                                                                                                                Submitted (NC
EMBL; AF32130
HSSP; P11362;
MGD; MGI:2150
GO; GO:000588
GO; GO:000551
GO; GO:000828
InterPro; IPR
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Mammalia; Eut
NCBI_TaxID=10
                                                                                  Receptor;
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SMART;
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(NOV-2000) to the
21301; AAL06296.1;
                                                                                         Signal.
                                                                                                                       150920; Fgfrll.
5886; C:plasma membrane; IDA.
5515; F:protein binding; IPI.
8285; P:negative regulation of cipr007110; Ig-like.
IPR003598; Ig_c2.
                                                                                                                                                                                                                                                                                                                                               1 (TrEMBLrel. 19, Created)
1 (TrEMBLrel. 19, Last sequence update)
4 (TrEMBLrel. 26, Last annotation update)
9 growth factor receptor 5 gamma precursor.
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1408; IGc2; 2.
150835; IG_LIKE; 2.
                        60.3%; ilarity 65.1%; Conservative
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theria; Rodentia; Sciurognathi; Muridae; Murinae; Mu
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438
                                                           46980 MW;
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; Mismatches
                          Score 1035.5; DB 2;
Pred. No. 7.7e-72;
Mismatches 14;
                                                                 gamma.
                                                                           Potential. fibroblast
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                                                            768D7E96126DD8AE CRC64;
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SORRED DRAFT RAPARA RAP
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C TISSUB=Kidney;
X MEDLINE=22388257; PubMed=12477932;
A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Strausberg R.L., Feingold E.A., Grouse L.H., Schuler G.D.,
A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
A Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
A Hopkins R.F., Jordan H., Moore T., Rubin G.M., Hong L.,
A Hopkins R.F., Jordan H., Bonaldo M.F., Casavant T.L., Scheetz T.E.
A Stapleton M., Joquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.
B Brownstein M.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.
A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A RA Rodriguez A.C., Grimwood J., Scheutz J., Myers R.M., Butterfield Y.
A RA Rodriguez A.C., Grimwood J., Scheutz J., Myers R.M., Butterfield Y.
A RA Rodriguez A.C., Grimwood J., Scheutz J., Myers R.M., Butterfield Y.
A RA Rodriguez A.C., Grimwood J., Scheutz J., Myers R.M., Butterfield Y.
B Grimwood J., Scheutz J., Myers R.M., Butterfield Y.
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B Grimwood J., Scheutz J., Myers R.M., But
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Proc. Natl. Acad. Sci. U.S.A [2]

SEQUENCE FROM N.A.

TISSUE=Kidney;

Strausberg R.;

Strausberg R.;

Submitted (SEP-2001) to the EMBL; BC013955; AAH13955.1;

InterPro; IPR003599; Ig.

InterPro; IPR007110; Ig-like InterPro; IPR003598; Ig_c2.

Pfam; PF00047; ig; 2.

SMART; SM00409; IG; 2.

SMART; SM00408; IGc2; 2.

PROSITE; PS50835; IG_LIKE; 2

SEQUENCE 350 AA; 37962 MW
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Q6PJN1;
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05-JUL-2004
05-JUL-2004
FGFRL1 prote
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Mammalia; Eutheria;
NCBI_TaxID=9606;
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
B Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Robards S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
Krzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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SEQUENCE FROM
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primata; Catarrhini; Hominidae;
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(SEP-2001) to the EMBL/GenBank/DDBJ databases
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350 AA; 37962 MW; C92A18DB4374A831 CRC64;
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B Ş Search completed: February Job time : 197 secs 2, 2005, 18:26:02

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